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如果她如果你看我的我的的话的话,我们就有这个人的话,我们就是我的人的话,我们就是我们的人的话,我们的人们的人,我们的人们的人,我们的人们的人们的人们的人们的人们的	
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Release 3.1A John F. Collins, Biocomputing Research Unit. Copyright (c) 1993-1998 University of Edinburgh, U.K. Distribution rights by Oxford Molecular Ltd

MPsrch\_pp protein - protein database search, using Smith-Waterman algorithm Tue Jan 18 09:51:49 2000; MasPar time 12.35 Seconds 702.420 Million cell updates/sec

Description: Perfect Score: Title: >US-09-332-063-2 (1-675) from US09332063.pep

rar output not generated.

Sequence: 4416

1 MPRAQPSSASYQPVPADPFA.....KTPIQILGQEPDAEMVEYLI 675

Searched: Scoring table: PAM 150 Gap 11

130303 seqs, 12848679 residues

Post-processing: Minimum Match 0% Listing first 45 summaries

Database: a-issued

1:5A\_COMB 2:5B\_COMB 3:PCT9\_COMB 4:backfiles1

Statistics: Mean 34.900; Variance 263.923; scale 0.132

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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14,	Sequence 14, Applicati	Sequence 2, Applicatio	sequence os, Applicati	, .	٠,		N,	Sequence 65, Applicati	32,	4	6, A	51,	Sequence 1, Applicatio	Sequence 1, Applicatio	2, A	19,	Sequence 52, Applicati	Sequence 51, Applicati	Sequence 54, Applicati	Sequence 2, Applicatio	Patent No. 5482709.	č	52720	Sequence 4. Application	Description	
	8.76e-04	4.90e-04	2.16e-04			2 430-04		9.54e-05	9.54e-05	5.96e-05	4.71e-05	5.96e-05	5.96e-05	5.96e-05	4.19e-05	3.31e-05	2.61e-05	2.61e-05	2.32e-05	6.32e-06	4.99e-06	4.996-06		2 536-07	Pred. No.	

## ALIGNMENTS

ADDRESSEE: Qu STREET: P.O. CITY: Madison	NUMBER OF SEQUENCE A	APPLICANT: John	GENERAL INFORMATION	Sequence 4, Applica	Sequence 4, Applicat		XXXXXX	US-08-217-327-4
Harles and Brady Box 2113	ES:	× 10	ON:	ation US/08217327	tion US/08217327	•		STANDARD; PI
	1 Protei			7				PRT; 21
	ns					•		214 AA.
	in							\$
	Cotton							
	Fiber							
	ADDRESSEE: Quarles and Brady STREET: P.O. Box 2113 CITY: Madison	ES es	APPLICANT: John, Maliyakal E APPLICANT: Barton, Kenneth A TITLE OF INVENTION: Immobilized Proteins in Cotton Fiber NUMBER OF SEQUENCES: 16 CORRESPONDENCE ADDRESS: ADDRESSE: Quarles and Brady STREET: P.O. Box 2113 CITY: Madison	NTO MATION: NT: John, Mal NT: Barton, K F INVENTION: OF SEQUENCES: ONDENCE ADDRES SSEE: Quarles T: P.O. Box 2 Madison	Mal Mal KES: S: RES	Sequence 4, Application US/08217327  Sequence 4, Application US/08217327  Patent No. 5474925  GENERAL INFORMATION: APPLICANT: Barton, Kenneth A  TITLE OF INVENTION: Immobilized Proteins in Cotton Fiber  NUMBER OF SEQUENCES: 16  CORRESPONDENCE ADDRESS: ADDRESSE: Quarles and Brady  STREET: P.O. Box 2113  CITY: Madison	Sequence 4, Application US/08217327  Sequence 4, Application US/08217327  Patent No. 5474925  GENERAL INFORMATION:  APPLICANT: John, Maliyakal E APPLICANT: Barton, Kenneth A  TITLE OF INVENTION: Immobilized Proteins in Cotton Fiber  NUMBER OF SEQUENCES: 16  CORRESPONDENCE ADDRESS:  ADDRESSEE: 0.0 Box 2113  CITY: Madison	Sequence 4, Application US/08217327  Sequence 4, Application US/08217327  Patent NO. 5474925  GENERAL INFORMATION: APPLICANT: John, Maliyakal E APPLICANT: Barton, Kenneth A TITLE OF INVENTION: Immobilized Proteins in Cotton Fiber NUMBER OF SEQUENCES: 16  CORRESPONDENCE ADDRESS: ADDRESSE: Quarles and Brady STREET: P.O. Box 2113 CITY: Madison

COUNTRY: USA.
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/217,327
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/812,233
FILING DATE: 19-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 1122990831
TELECOMMUNICATION: TELEPHONE: 608-251-5000
TELEPHONE: 608-251-9166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 214 amino acids
TYPE: amino acid
TOPOLOGY: 11 near 1122990831

TYPE: ami

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01-JAN-1900
                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                         SPOROZOITE 21.5 KB ANTIGEN, AC-6B
NUMBER OF SEQUENCES: 11
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: JACOBSON, JAMES W.; STRAUSBERG, SUSAN L.; STRAUSBERG, SUSAN L.;
                       XXXXXX
                                              5482709-6
                                                                                                                                                                                                                                                                                                          SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AUGUSTINE, PATRICIA C.; DANFORTH, HARRY D.
TITLE OF INVENTION: GENETICALLY ENGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
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SEQUENCE 214 AA;
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                                                                                                                                                         81 ATGAAAAPAAATAAAGAAGAAKAAEARTETEGAAGAAEAEKAKTQAATAATTAAAARAA 140
                                                                                                                                                                                                   Local Similarity
les 56; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       32 STPATFTPPASTPPP-TTQAPPTP-TATPPPVSTPPPTSSPPPVTASPPPVSTPPPSSP-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/V
FILING DATE: 12-SEP-1990
PRIOR APPLICATION DATA:
                                                                                        VPAPAAAQASAPAQTQAPTSAPAVAPTPAPTPTPAVAQAE 617
                                                                                                                                   PSPATAAATAAAVSPAAAGQIPAAASVASAAAVAPSAAAAAAVQVAPAAPAPVPAPALVP 577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQASAPAQTQAPTSAPAVAPTPAPTPTPAVAQAEVPASPATGPGPHRLSIPSL 636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAATAAAVSPAAAGQIPAAASVASAAAVAPSAAAAAAVQVAPAAPAPVPAPALVPVPAPA
                                                                                                                                                                                                                                                                                                                             FILING DATE: 05-JUL-1988
APPLICATION NUMBER: 746,520
FILING DATE: 19-JUN-1985
APPLICATION NUMBER: 627,811
                                                                                                                                                                                                                                                                                                               FILING DATE: 05-JUL-1984
                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 215,162
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20649 MW; 275674 CN;
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                                           PRT;
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Pred. No. 4.99e-06;
51; Mismatches 53
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Pred. No. 2.53e-07
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FILING DATE: 05-JUL-1989
APPLICATION NUMBER: 746,520
FILING DATE: 19-JUN-1985
APPLICATION NUMBER: 627,811
FILING DATE: 05-JUL-1984
SEQ ID NO:6:
                                                                                                                                                                                                                                                                                                                         XXXXXX
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TITLE OF INVENTION: EIMERIA ANTIGENIC COMPOSITION WHICH
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                                                                                                                                                                                                                                                                    Sequence 2, Application US/08450360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   458
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                                                                                                                                                                                                                                                                                                                                                                                               578 VPAPAAAQASAPAQTQAPTSAPAVAPTPAPTPTPAVAQAE 617
                                                                                                                                                                                                                                    Sequence 2, Application US/08450360 Patent No. 5656457
                                                                                                                                                                                                                                                                                                                                                                                                                      141 AASAAGKPGGHAAASTAAKAAQEKAAKAANAATAATASKE 180
                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.9%;
Local Similarity 35.0%;
les 56; Conservation
                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81 ATGAAAAAAAAAAAGAAGAAKAAEARTETEGAAGAAAEAEKAKTQAATAATTAAAARAA 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                             APPLICANT: Parkes, APPLICANT: Coates,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NÜMBER: US/08/
FILING DATE: 08-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 581,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 10 CURRENT APPLICATION DATA:
                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                            PSPATAAATAAAVSPAAAGQIPAAASVASAAAVAPSAAAAAAVQVAPAAPAPVPAPALVP
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                                                                       ZIP:
                                                                                  COUNTRY:
                                                                                               STATE:
                                                                                                             CITY:
                                                                                                                        STREET:
                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 12-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 180
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                                                                                             Tiburon
California
                                                                                                                     6 Mariposa Court
                                                                                 USA
                                                                                                                                                                                                          Parkes, Deborah Lynn
         PatentIn Release #1.0,
                                                                                                                                   Leona L.
                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                                                                                                                                                 Stephen Ralph
Herpes Simplex Virus Type 2-Glycoprotein G
Proteins and Polypeptides
                                                                                                                                   Lauder
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 259; DB 4;
Pred. No. 4.99e-06;
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         Version #1.30 (EPO)
                                                                                                                                                                                                                                                                                                                                              486 AA
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Best Local :
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                                                                                                                                                                                                                                                                                              XXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                         Sequence 54, Application US/08557309B Patent No. 5916572
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                                                                                                                                                                                                                                                                                                                                                                                                                                                318 SSDSPTSAPEKTPLPVSATAMAPSVDPSAEP-TAPATTTPPDEMATQAA-TVAVTPE-ET 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                     COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC -DOS/MS-DOS
                    SOFTWARE: Patentin Release -#1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                             APPLICANT: Reed, Steven G.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Lodes, Michael J.
                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein
JENCE 486 AA; 50029 MW; 1269874 CN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Lauder, Leona L.
REGISTRATION NUMBER: 30,863
REFERENCE/DOCKET NUMBER: D-0
TELECOMMUNICATION INFORMAÇION:
TELEPHONE: 415-435-2034
                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                          APPLICANT:
                                                                                                     STREET: COLUMN STREET: Seattle
                                                                                                                                                                                                                                                                                                                                                                                        aaataaavspaaagqipaaasvasaaavapsaaaaavqvapaapapvpapalvpvpapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                  AAQASAPAQTQAPTSAPAVAPTPAPTPTPAVAQAEVPASPATGPGPHRLSIPSLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/450,360 FILING DATE: 25-MAY-1995
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                                                                                                                                                                                                               INFORMATION:
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               CATION NUMBER:
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ilarity 27.4%;
Conservative
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             US/08/557,309B
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Pred. No. 6.32e-06;
71; Mismatches 50
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      REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: A55556-3/BIR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8700
                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                         ZIP:
                                                                             FILING DATE:
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(415) 494-8771
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596 TSAPAVAPTPAPTPTPAVAQAEVPASPATGPGP 628
                                176 АКАЛАРРАКАЛАРРАКАЛАРРАКАЛАРРАЕЛАР 208
                                                              536 GQIPAAASVASAAAVAPSAAAAAVQVAPAAPAPVPAPALVPVPAPAAAQASAPAQTQAP 595
                                                                                              116 ААРРАКАААРРАКТААРРАКТААРРАКАААРРАКАААРРАКАААРРАКААААРАКААААР 175
                                                                                                                                 476 APVAAAATAAAITATAATITTIMVAAAPVAVAAAAAPAAAAAPSPATAAATAAAVSPAAA 535
                                                                                                                                                   TELEFAX: (206) 682-60 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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CE 219 AA; 21347 MW; 161993 CN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Maki, David J.
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Similarity 37.9%;
58; Conservative
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Pred. No. 2.32e-05;
Mismatches 52;
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Sequence 51, Application US/08642255 STANDARD; PRT; 234 A

Patent No. 5//January Patent No. 5//January Patent INFORMATION: TITLE OF INVENTION: APPLICANT: CAPPELLO, Joseph APPLICANT: FERRARI, Franco A. 51, Application US/08642255 o. 5773249 Protein Polymers High Molecular Weight Collagen-Like

SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk ATTORNEY/AGENT INFORMATION: CLASSIFICATION: 435 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS APPLICATION NUMBER: 94111-4187 ROWLAND, Bertram California E: FLEHR, HOHBACH, TEST, ALI 4 Embarcadero Center, Suite Francisco US/08/642,255 ALBRITTON Version #1.30

& HERBERT

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RESULT
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nes 51; Conser
         INFORMATION
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                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                   ATTORNEY/AGENT INFORMATION: NAME: ROWLAND, Bertram I
                                                                                                                                                                                                                                                                                                APPLICANT: FERRARI, Franco A.
TITLE OF INVENTION: High Molecular Weight Collagen-Like
TITLE OF INVENTION: Protein Polymers
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MOLECULE TYPE:
TELEPAC
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TELEX: 910 27
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                                                  REFERENCE/DOCKET NUMBER: A5 ELECOMMUNICATION INFORMATION:
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                                                                                                           CLASSIFICATION:
                                                                                                                       FILING DATE:
                                                                                                                                                                                                                        COUNTRY:
                                                                          REGISTRATION NUMBER:
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234 AA; 18406 MW;
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                IE: (415) 494-870
(415) 494-8771
910 277299 FHT UF
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4 Embarcadero Center, Suite 3400
an Francisco
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Pred. No. 2.61e-05;
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APPLICANT: Spaete, Richard and Jack
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          TELEX: 380816 CooleyPA INFORMATION FOR SEQ ID NO:
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Local Similarity 24.8%;
hes 51; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDENNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
DUENCE 561 AA; 46438 MW; 1
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STREET: 9alo Alto
CITY: Palo Alto
CTATE: California
                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: April 1
                                                                                                                                                                                                             MEDIUM TYPE: floppy disk
COMPUTER: IMP PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0
CURRENT APPLICATION DATA:
SEQUENCE CHARACTERISTICS
                                            REFERENCE/DOCKET NUMBER: AV TELECOMMUNICATION INFORMATION: TELEPHONE: 415-843-5163
                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Luann Cserr
REGISTRATION NUMBER: 31,
                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                   TELEFAX:
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5 Palo Alto Square
                                   415-857-0663
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Non Splicing Variants of gp350/220
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                                                                                                                                     April 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
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                                                                                  31,822
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Pred. No. 2.61e-05;
84; Mismatches 63
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Length 2441; Indels 10;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/08194468 Patent No. 5750336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          549
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               530 PTPNATSPTPAVTTPTP-NATSPTLGKTSPTSAVTTPTPNATSPTL-GKTSPTSAVTTPT 587
                                                                                                                         TELEFAX: (619)-546-93 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: au....
TOPOLOGY: linear
MOLECULE TYPE: protein
MOLECULE TYPE: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/194,468 FILING DATE: 10-FEB-1994
                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (619)-546-4737
                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PTPAVAQAEVP-ASPATGPGPHRLSIPSLTCNPDKTDGPVFH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 444 CITY: Los Angeles
                                                                                                                                                                                                                  NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
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                                                                               ENGTH:
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                                                     amino acid
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                                                                               2441 amino acids
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MORTHIDY, MATC R.

VENTION: ASSAYS FOR THE IDENTIFICATION OF

VENTION: COMPOUNDS WHICH INHIBIT ACTIVATION OF CAMP AND MITOGEN
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                                                                                                                                              (619)-546-9392
                                                                                                                                                                                                                                                                                                                                                                                                                   IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Floppy disk
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Pred. No. 3.31e-05;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          944 QSSQQQPTPVHTQPPGTPLSQAAASIDNRVPTPSTVTSAETSSQQPGPDVPMLEMKTEVQ 1003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         431 TPSPVPPSTPLLSAHSKTGSRDCSTQTERGTESNKTAAVAPISV-PAPVAAAATAAAITA
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                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Haughton, Raymond
APPLICANT: Skeiky, Yasir A.W.
TITLE OF INVENTION: COMPOUNDS
TITLE OF INVENTION: OF T. CRU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                       REGISTRATION NUMBER:
                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE:
COPOLOGY:
                                          ENGTH:
                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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Similarity 23.4%;
58; Conservati
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                                                                                                              (206)
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                                                                                                                                                                                                                       David J.
                                                                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                                                                                                                                                             15-SEP-1997
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                                                                                                         682-6031
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Pred. No. 4.19e-05;
89; Mismatches 91
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                                                                                                                                                                             210121.406C1
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                                 Query Match
Best Local Similarity 37.5%,
Conservative
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SEQUENCE 262 AA;
104 RERARREDAAAAAAKQKAAAAKKAAAAAPSGKKSAKAAIAPAKAAAAAPAKAAAAAP-AKAAAA 162
                                                                                                                                               SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 571 PAPALVPVPAPAAAQASAPAQTQAPTSAPAVAP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              511 APAAAAAPSPATAAATAAAVSPAAAGQIPAAASVASAAAVAPSAAAAAAAVQVAPAAPAPV 570
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  163 РАКАЛААРАКАЛААРАКААТАРАКААЛАРАКТАЛАРАКАЛАРАКАЛАРАКАЛТАРАКАЛ 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         451 RDCSTQTERGTESNKTAAVAPISVPAPVAAAATAAAITATAATITTTMVAAAPVAVAAAA 510
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                                                                                                                                                                                                                                                              TELEFAX: (206) 682-60 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
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nes 58; Conser
                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 262 amino acid
                                                                                                                                                              TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 98104-, COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                        ELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                NAME: Maki, David J. REGISTRATION NUMBER: 31,392
                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                       JE TYPE: protein 262 AA; 27103 MW; 216181 CN;
                                                                                                                                                                                                   amino acid
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                                                                                                                                                                                                                    262 amino acids
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                                                                               5.48;
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                                                                                                                                                                                                                                                                                                   206) 622-4900
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37.98;
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27103 MW; 216181 CN;
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NO: 1:
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                                                                               Score 238; DB 1;
Pred. No. 5.96e-05
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Pred. No. 5.96e-05
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                                                          Mismatches
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Best Local Similarity 37.9%;
                                                                                                                                                                                      SEQUENCE
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                      511 APAAAAAPSPATAAATAAAVSPAAAGQIPAAASVASAAAVAPSAAAAAAVQVAPAAPAPV 570
                                                164 РАКАЛААРАКАЛАРАКАЛТАРАКАЛАРАКТАЛАРАКАЛАРАКАЛАРАКАЛТАРАКАЛ 223
                                                                          451 RDCSTQTERGTESNKTAAVAPISVPAPVAAAATAAAITATAATITTTMVAAAPVAVAAA 510
                                                                                                  105 RERARREDAAAAAAAKQKAAAKKAAAPSGKKSAKAAIAPAKAAAAPAKAAAAP-AKAAAA 163
224 AAPA-KAATAPAKA-ATAPAKAAAAPAKAATAP 254
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 451 RDCSTQTERGTESNKTAAVAPISVPAPVAAAATAAAITATAATITTTMVAAAPVAVAAAA 510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
                                                                                                                                                                                                                                                   SEQUENCE CHARACT
                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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                                                                                                                                                                                                 TYPE: amino
STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 6300 of CITY: Seattle
                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                          TELEPHONE:
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                                                                                                                                                                                  263 AA; 27231 MW; 218279 CN;
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                                                                                                                                                                                                                           amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                          IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                           Score 238; DB 2;
Pred. No. 5.96e-05;
                                                                                                                                Mismatches
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Best Local
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                                      586 ASAPAQTQAPTSAPAV-APTPA-PTPTPAV-AQAEVPASPATGPGP-HRLSIPSLTCNPD
                                                                                                           470 APISVPAPVAAAATAAAITATAATITTTMVAAAPVAVAAAAAPAAAAAP-SPATAAATAA 528
      642 KTDGPVF 648
                       216 TTPAPPY 222
                                                            156
                                                                        529 AVSPAAAGQIPAAASVASAAAVAPSA-AAAAAVQV-APAAPAPVPAPAL-VPVPAPAAAQ
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                                                                                                                                               5.48;
Local Similarity 28.98;
hes 54; Conservation
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                                                                                          96 ҮКРРАРАРРТКАРТРРҮКРРАРАРРТКАРТРРҮКРРАРАРРТКАРТРРЕКРРАРАРРТКА 155
                                                                                                                            36 АРУРУКАРІРАРРУКРРТТРУРРУКАРТРАРРТКСРТРУКРРТКАРТРРУКРРТКАРТРР 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6, Application US/08580545B
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                     MOLECULE TYPĖ: protein
JENCE 297 AA; 31270 MW; 517269 CN;
                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 297 amino acids
                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Bretschneider, Barry E.
REGISTRATION NUMBER: 28,055
                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy
                                               PTPPYKPPTPAPAPPVKAPTPPYMPPTPPTKAPTPAPAPPTKAPTPPYKPPVPTPPVKPP 215
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CORRESPONDENCE ADDRESS:
Fish & Richardson P.C.
ANDRESSEE: Fish treenth Street, NW
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                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
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Susumu, Nishiguchi
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                                                                                                                                                                                                          linear
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                                                                                                                                                 Score 240; DB 2;
Pred. No. 4.71e-05;
66; Mismatches 59
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Best Local Similarity 28.6%;
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                            398 TPPTTPAPTTPPPTSTHATPRPTTPGPQT-TPPGPAT-PGP 436
588 APAQTQAPTSAPAVAPTPAPTPTPAVAQAEVPASPATGPGP
                                                             528 AAVSPAAAGQIPAAASVASAAAVAPSAAAAAAVQVAPAAPAPVPAPALVPVPAPAAAQAS
                                                                                         339 TTPPDEMATQ-AATVAVTPEETAVASPPATASVESSPLPAAAAATPGAGHTNTSSASAAK 397
                                                                                                                          468 AVAPISVPAPVAAAATAAAITATAATITTTMVAAAPVAVAAAAAPAAAAAASAAAASAATAA 527
                                                                                                                                                        281 GVDP-DVRAPRTGRRLMALTEDTSSDSPTSAPEKTPLPVSATAM-APSVDPSAEPTAPAT 338
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                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 447 amino acids
TYPE: amino acid
TOPOLOGY: linear
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: CIBUTON
                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Lauder, Leona L.
REGISTRATION NUMBER: 30,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC-
OPERATING SYSTEM:
SOFTWARE: Patentl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/01
FILING DATE: 25-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  California
                                                                                                                                                                                                                                                                                                                                                                                                415-435-0727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 Mariposa Court
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
Patentin Release #1.0, Version #1.30 (EPO)
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Stephen Ralph
Herpes Simplex Virus Type 2-Glycoprotein G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US 08/129,021
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                                                                                                                                                                                            Score 238; DB 1; Le
Pred. No. 5.96e-05;
57; Mismatches 53;
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US-08-642-255-32

STANDARD;

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st Local Similarity 22.1%;
matches 49; Conservative
                                                                                                                                                                                                                                                                                   SEQUENCE
606 APTPTPAVAQAEVPASPATGP-GPHRLSIPSLTCNPDKTDGP 646
                                                                                                                                                              213 PGPPGPPGPAGPVGSPGAPGPPGPPGPPGPPGAPGPPGPPGP 254
                                                  546 SAAAYAPSAAAAAYQVAPAAPAPVPAPALVPYPAPAAAQASAPAQTQAPTSAPAYAPTP 605
                                                                                                         487 ITATAATITTTMVAAAPVAVAAAAAPAAAAAP-SPATAAATAAAVSPAAAGQIPAAASVA 545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 32, Application US/08642255
                                                                           155 PPGPPGPPGAPGPPGPPGPPGPPGPPGPA-GPVGSPGAPGPPGPPGPPGAPG-PPGP 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 5773249
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 32, Application US/08642255 Patent No. 5773249
                                                                                                                                                                                           TELEFAX: (415) 494-8771
TELEX: 910 277299 FHT UR
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8770
TELEFAX: (415) 494-8771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                             LENGTH: 330 amino
TYPE: amino acid
STRANDEDNESS: sing
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 20, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                JLE TYPE: protein
330 AA; 28876 MW; 625100 CN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
1: 4 Embarcadero Center, Suite 3400
San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ROWLAND, Bertram I
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                                                                                                                                                                                                                                                                                                                                                   330 amino acids
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WENTION: High Molecular Weight Collagen-Like
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/08/642,255
                                                                                                                                                                                                                        Score 234; DB 1;
Pred. No. 9.54e-05;
90; Mismatches 76
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                                                                                                                                                                                                                        Indels
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Search completed: Tue Jan 18 09:52:05 2000 Job time: 16 secs.

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Release 3.1A John F. Collins, Biocomputing Research Unit. Copyright (c) 1993-1998 University of Edinburgh, U.K. Distribution rights by Oxford Molecular Ltd

Run on: MPsrch\_pp protein - protein database search, using Smith-Waterman algorithm Tue Jan 18 09:50:44 2000; MasPar time 43.38 Seconds 849.350 Million cell updates/sec

Description: Perfect Score: Sequence: Title: >US-09-332-063-2 (1-675) from US09332063.pep 4416

Jular output not generated.

1 MPRAQPSSASYQPVPADPFA.....KTPIQILGQEPDAEMVEYLI 675

Scoring table: PAM 150 Gap 11

Searched: 179066 seqs, 54579741 residues

Post-processing: Minimum Match 0% Listing first 45 summaries

Database: sptremb19

1:sp\_archea 2:sp\_bacteria 3:sp\_fungi 4:sp\_human 5:sp\_invertebrate 6:sp\_mammal 7:sp\_mhc 8:sp\_organelle 9:sp\_phage 10:sp\_plant 11:sp\_rodent 12:sp\_unclassified 13:sp\_vertebrate 14:sp\_virus

S'wilstics: Mean 54.237; Variance 271.816; scale 0.200

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

1100 887 665 4 4 3 3 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Result
372 365 345 347 347 340 340 335 335 337 337 327 327 327 327 327 327 327 327	Score
6.66.89991944	Query Match
507 3534 807 867 2378 788 60 2378 861 216 216 216 216 270 851 750 770 825 851 473 1188 871 881 781 871 871 871 871 871 87	Length
114 114 114 114 114 114 114 114 114 114	DB
0.130.28 0.39266 0.39266 0.39781 0.39782 991365 991365 0.18510 0.18510 0.23635 0.23635 0.228501 0.24888 0.42854 0.42855 0.42854 0.4285	ID
ANTIFREZE GLYCOPEPTID COUNTERPART OF HSV-1 G INSECT INTESTINAL MUCI MEMBRANE GLYCOPROTEIN. CODED FOR BY C. ELEGAN INSECT INTESTINAL MUCI MEMBRANE GLYCOPROTEIN. SIMILAR TO LONG TANDEM MUCIN (FRAGMENT). MYCIN (FRAGMENT). HYPOTHETICAL 170.5 KD POSITIONAL COUNTERPART F56H9.1 PROTEIN. C12D12.1 PROTEIN. C12D12.1 PROTEIN. F55B11.3 PROTEIN. F55B11.3 PROTEIN. (VSP-3) PRECURSOR. EXTENSIN-LIKE PROTEIN HYPOTHETICAL 74.5 KD P F47B8.5 PROTEIN.	Description
1.18e-15 3.96e-15 3.96e-14 1.12e-14 2.47e-13 2.93e-13 6.88e-13 1.15e-12 2.69e-12 2.69e-12 2.7e-12 2.7e-12 3.41e-11 7.92e-11 9.37e-11 1.31e-10 2.56e-10 0.75e-10 1.36e-10 1.36e-10 1.36e-10	Pred. No.

4.5	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21
259	261	262	259	263	264	266	270	269	271	270	268	278	280	279	278	280	278	282	283	284	287	285	288	288
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•	-	L Q62605		2 Q44562		) Q43558		_		_		075112			5 Q29070					_	•	5 Q290 <b>71</b>		1 Q14883
SIMILAR TO A. FAECALIS	MEGAKARYOCYTE STIMULAT	APOMUCIN PRECURSOR.	SMALL INTESTINAL MUCIN	SIALIDASE.	AFGP POLYPROTEIN PRECU	PROLINE RICH PROTEIN P	HIGH MOLECULAR MASS NU	EXTENSIN (FRAGMENT).	MUCIN (FRAGMENT).				AFGP POLYPROTEIN PRECU	GLYCOPROTEIN 150.	GASTRIC MUCIN (FRAGMEN	PUTATIVE PROLINE-RICH	н	N)		CEL	a	GASTRIC MUCIN (FRAGMEN	_	INTESTINAL MUCIN (FRAG
2.23e-07	1.61e-07	1.37e-07	2.23e-07			7.12e-08	.70e		3.14e-08		5.13e-08	9.92e-09								3.68e-09		3.12e-09	1.90e-09	1.90e-09

## ALIGNMENTS

рь Оу	Qy Db	рb	Qy Qy	Ma Be	SPITE	RA RA RT RT	RESOLT AC OD DT OD OT OC
309 ATPATPATAATAATAATAATAATAAAAPAATAATPATA 350 	250 TAATPARAARAATPATAATLATAATPATPATAATDATAATDARAATPAT-PATA 308 	190 TPATAATAATAATAATAATAATAATAATPARAATAATPATAPTPATAATPATAATAATAATA 249 	130 RAATPATAATPATAATAATAATAATAATAARAATPATAATAATAATAATAARAATPATAARAARAA 189    ::::  ::   :  ::  ::  ::  ::   :   :	Query Match  8.4%; Score 372; DB 13; Length 507; Best Local Similarity 33.3%; Pred. No. 1.18e-15; Matches 74; Conservative 74; Mismatches 71; Indels 3; Gaps 3;		SEQUENCE FROM N.A.  MEDLINE; 97268653.  CHEN L., DEVRIES A.L., CHENG C.H.C.;  "Convergent evolution of antifreeze glycoproteins in Antarctic notothenioid fish and Arctic cod.";  PROC. NATL. ACAD. SCI. U.S.A. 94:3817-3822(1997).  EMBL; 043200; G2078483;	O13028 PRELIMINARY; PRT; 507 AA. O13028; PRELIMINARY; PRT; 507 AA. O13028; O1, JUL-1997 (TREMBLREL: 04, CREATED) O1-JUL-1997 (TREMBLEEL: 04, LAST SEQUENCE UPDATE) O1-NOV-1998 (TREMBLEEL: 08, LAST ANNOTATION UPDATE) ANTIFREEZE GLYCOPEPTIDE AFGP POLYPROTEIN PRECURSOR. BOREOGADUS SAIDA. EUKARYOTA; METRAZOA; CHORDATA; VERTEBRATA; ACTINOPTERYGII; NEOPTERYGII; FELEOSTEI; EUTELEOSTEI; PARACANTHOPTERYGII; GADIFORMES; GADOIDEI; GADIDAE; BOREOGADUS. [1]

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Best Local S
Matches 8
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STRAIN-NS80567;

( MEDLINE; 98264497.

A TELFORD E.A.R., WATSON M.S., PERRY J., CULLING THE DNA sequence of equine herpesvirus-4.";

The DNA sequence of equine herpesvirus-4.";
                                                                                                                                                                 MEDLINE;
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01-JAN-1998
01-NOV-1998
                                                                                                                                                                                                                                                                                                                                                    01-JAN-1998 (TREMBLREL 05, CREATED)
01-JAN-1998 (TREMBLREL 05, LAST SEQUENCE UPDATE)
01-OV-1998 (TREMBLREL 08, LAST ANNOTATION UPDATE)
INSECT INTESTINAL MUCIN IIM22
TRICHOPLUSIA NI (CABBAGE LOOPER)
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018511;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  039266;
   SEQUENCE
                                            EMBL;
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                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                       PTERYGOTA;
                                                                                                                                                                                                                                                                                                                                   EUKARYOTA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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                                                                                                              "Molecular cloning and 
nucin cDNA.";
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VIRUSES; DSDNA VIRUSES,
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                                                                        BIOL.
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                                         AF000606;
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80; Conser
                                                          CHEM. 272:16663-16669(1997).
                                                                                                                                                                 97341213.
GRANADOS R.R.;
                                                                                                                                                                                                                                                                                                LEPIDOPTERA; NOCTUOIDEA;
                                                                                                                                                                                                                                                                                                                                   METAZOA; ARTHROPODA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3534 AA;
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(TREMBLREL. 05, LAST SEQUENCE UP
(TREMBLREL. 08, LAST ANNOTATION
OF HSV-1 GENE UL36 AND VZV GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8.3%;
llarity 32.3%;
Conservative
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      AA;
                                     G2224921
      83408
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VARICELLOVIRUS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       379957
                                                                                                                               sequencing
   MW.
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83; 1
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LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 365; DB 14;
Pred. No. 3.96e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
15E08C34 CRC32;
                                                                                                                                                                                                                                                                                             TRACHEATA; HEXAPODA;
DEA; NOCTUIDAE; TRICHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CE599A95 CRC32;
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                                                                                                                                  novel
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                                                                                                                               invertebrate
                                                                                                                                                                                                                                                                                             APODA; INSECTA;
TRICHOPLUSIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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                                                                                                                               intestinal
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RESULT
ID P9
AC P9
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OS C4
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Best Local S
Matches 5
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Best Local
                                                    01-MAY-1997 (
01-MAY-1997 (
01-NOV-1998 (
CODED FOR BY
                                                                                                          P91365
RHABDITINA;
                     CAENORHABDITIS ELEGANS. EUKARYOTA; METAZOA; NEM
                                            K06A9.1.
                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-BK343;
KIRISAWA R., KOBAYASHI T., KA
J. EQUINE SCI. 7:79-87(1996).
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01-JAN-1998 (TREMBLREL.
01-JAN-1998 (TREMBLREL.
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                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EQUINE HERPESVIRUS 1. VIRUSES; DSDNA VIRUSES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALPHAHERPESVIRINAE; VARICELLOVIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEMBRANE GLYCOPROTEIN.
                                                                                                                                                      609
                                                                                                                                                                         348
                                                                                                                                                                                                                     288
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                                                                                                                                                                                                                                                         228 ATTTAATTTAATTTAATTTAATTTAATTSSATTAATTTAATTTAATTTAATTTAATTTAA 287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            579
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                                                                                                                                                PTPAVAQAEVPASPATGPGPHRLSIPSLTCNPDKTDG
                                                                                                                                                                       PSASTATSATPTSTSTSAAATT-STPTPTSAATSAES 383
                                                                                                                                                                                                               TTTAATTTAATTTAATTTAATTTAATTTAATTTAATTTAATTTGSPTSGSTSTTGASTST
                                                                                                                                                                                                                                      TAATITTTMVAAAPVAVAAAAAPAAAAAPSPATAAATAAAVSPAAAGQIPAAASVASAAA
                                                                                                                                                                                                                                                                               STPSPVPPSTPLLSAHSKTGSRDCSTQTERGTESNKTAAVAPISVPAPVAAAATAAAITA
                                                                                                                                                                                                                                                                                             VAPSAAAAAAVQVAPAAPAPVPAPALVPVPAPAAAQASAPAQTQAPTSAPAVAP-TPAPT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TATAA-TITTTMV-AAAPVAVAAAAAPA-AAAAPSPATAAATAAAVSPAAAGQIPAAASV
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                                                                                                                                                                                                                                                                                                                             h 7.7%;
Similarity 26.3%;
57; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 8.1%;
Similarity 34.2%;
                                                   (TREMBLREL.)
(TREMBLREL.)
(TREMBLREL.)
(TREMBLREL.)
           RHABDITOIDEA;
                                                                                                                                                                                                                                                                                                                                                                         867 AA;
                                                                                                           PRELIMINARY;
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                    NEMATODA;
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CDNA
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       DDA; SECERNENTEA; RHABDITI
RHABDITIDAE; PELODERINAE;
                                                  LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDAT
A YK50C7.5.
                                                                                                                                                                                                                                                                                                                                                                                                                   KAWAKAMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA STAGE;
                                                                                                                                                                                                                                                                                                                           Score 341; DB 14;
Pred. No. 2.47e-13;
91; Mismatches 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CREATED)
                                                                                    CREATED)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDATE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 359; DB 5;
Pred. No. 1.12e-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
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                   RHABDITIA;
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                                                                                                                                                                                                                                                                                                                             Indels
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       CAENORHABDITIS.
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RESULT
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RN EN SI
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Best Local
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                                                                                                      01-JAN 1998 (TREMBLREL 05, CREATED)
01-JAN-1998 (TREMBLREL 05, LAST SEQUI
01-NOV-1998 (TREMBLREL 08, LAST ANNO
INSECT INTESTINAL MUCIN IIM14
TRICHOPLUSIA NI (CABBAGE LOOPER),
EUKARYOTA; METAZOA; ARTHROPODA; TRACHI
PTERYGOTA; LEPIDOPTERA; NOCTUOIDEA; N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
STRAIN-BRISTOL N2;
GEISEL C., GATTUNG S.;
SUBMITTED (JAN-1997) T
                                                                                                                                                                                                        018510;
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BONFIELD J., BURTON J., CONNELL M., COPSEY T., COOPER J., COULSON A
CRAXTON M., DEAR S., DU Z., DURBIN R., FAVELLO A., FULTON L.,
GARDNER A., GREEN P., HAWKINS T., HILLIER L., JIER M., JOHNSTON L.,
JONES M., KERSHAW J., KIRSTEN J., LAISTER N., LATREILLE P.,
LIGHTNING J., LLOYD C., MCMURRAY A., MORTIMORE B., O'CALLAGHAN M.,
PARSONS J., PERCY C., RIFKEN L., ROOPRA A., SAUNDERS D., SHOWNKEEN
SMALDON N., SMITH A., SONNHAMMER E., STADEN R., SULSTON J.,
SMITH A., SONNHAMMER E., STADEN R., SULSTON J.
                        "Molecular cloning and mucin cDNA.";
                                                   SEQUENCE FROM N.A.
MEDLINE; 97341213.
WANG P., GRANADOS R.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; U80846; G1707247;
SEQUENCE 2378 AA; 225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WATERSTON R.;
SUBMITTED (APR-1997)
                                                                                                                                                                                                                                                                                                                                 548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUBMITTED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "2.2 Mb of contiguous nucleotide elegans.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                THIERRY-MIEG J., THOMAS K., VAUDIN M., WATSON A., WEINSTOCK L., WILKINSON-SPRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NATURE 368:32-38(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN-BRISTOL N2;
MEDLINE; 94150718
                                                                                                                                                                                                                                                                                                      406
                                                                                                                                                                                                                                                                                                                                                            346
                                                                                                                                                                                                                                                                                                                                                                                       489
                                                                                                                                                                                                                                                                                                                                                                                                                287
                                                                                                                                                                                                                                                                                                                                                                                                                                              430
                                                                                                                                                                                                                                                                                                                                                                                                                                                        227 PTTTTITPTTTVVPASTSTQPAPASSPTTQPAPATSPTTQPAPATSPTTQPAPATSPTTQ 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.7%;
Local Similarity 27.4%;
les 60; Conservative
              BIOL
                                                                                                                                                                                                                                                                       TPTPAVAQAEVPASPATGPGPHRLSIPSLTCNPDKTDGP
                                                                                                                                                                                                                                                                                                                                                     QPTPATSPTTQPTPATSPTTQPTPATSPTTQPAPATSPTTQPAPATSPTTQPAPATSPTT
                                                                                                                                                                                                                                                                                                 QPAPATSPTTQPA-PATSPTTQPAPATSPTTQPAPATSP 443
                                                                                                                                                                                                                                                                                                                             AAVAPSAAAAAVQVAPAAPAPVPAPALVPVPAPAAAQASAPAQTQAPTSAPAVAPTPAP
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                                                                                                                                                                                                                                                                                                                                                                                                                                     STPSPVPPSTPLLSAHSKT-GSRDCSTQTERGTESNKTAAVAPISVPAPVAAAATAAAIT 488
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                                                                                                                                                                                                                     PRELIMINARY;
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                                   sequencing
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                                                                                                                                                           CREATED)
LAST SEQUENCE UP
LAST ANNOTATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 340; DB 5;
Pred. No. 2.93e-13;
78; Mismatches 7.7
                                                                                                                                                                                                                    PRT;
                                                                                                      TRACHEATA; HEX
DEA; NOÇTUIDAE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    765EDB23 CRC32;
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f
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                                      novel
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                                                                                                                                                                                                                    A
                                                                                                                                                                          UPDATE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DATA
                                                                                                                    HEXAPODA;
                                   invertebrate intestinal
                                                                                                                                                                                                                                                                        646
                                                                                                       TRICHOPLUSIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 2378;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BANKS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                III of
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023635;
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SEQUENCE
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KIRISAWA R., KOBAYASHI T., KAN
J. EQUINE SCI. 7:79-87(1996).
EMBL; D88733; D1020845; -.
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VIRUSES; DSDNA VIRUSES,
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mes 55; Conser
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                                        J. 01, CREATED)
L. 01, LAST SEQUENCE UPDATE)
L. 08, LAST ANNOTATION UPDATE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 335; DB 14;
Pred. No. 6.88e-13;
90; Mismatches 71;
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LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDAT
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Pred. No. 6.88e-13;
77; Mismatches 58;
                                                                                                                                                                                    PRT;
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CAENORHABDITIS ELEGANS EUKARYOTA; METAZOA; NEI

NEMATODA;

SECERNENTEA;

RHABDITIA;

RHABDITIDA;

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01-NOV-1996 (TREMBLREL.
01-NOV-1998 (TREMBLREL.
                                                                                                                                                                 EMBL;
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STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                          MACACA MULATTA (RHESUS MACAQUE)
EUKARYOTA; METAZOA; CHORDATA; V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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BONFIELD J., BURTON :
                                                                                                               SEQUENCE
                                                                                                                                                                                                                                      "Expression of MUC2 gene
transcriptional level in
                                                                                                                                                                                                                                                                AN G., LUO G., WU R.;
"Expression of MUC2 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MUCIN (FRAGMENT)
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                                                                                                                                                                                                                                                                                                           MEDLINE; 94235322.
                                                                                                                                                                                                                                                                                                                                                ISSUE-TRACHEA
                                                                                                                                                                                                                                                                                                                                                                                                                    CATARRHINI; CERCOPITHECIDAE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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gans.":
                                                                                                                                                                                      J. RESPIR. CELL
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                                                                                                                                                              U00483;
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1 AA; 77123 MW;
                                                                                                                                                              G437055;
                                                                                                            Ä
                                                                                                                                                            MOL. BIOL.
                                  7.4%;
22.0%;
                                                                                                            21692 MW;
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                                                                                                                                                                                                                                is down-regulated by vitamin A at the vitro in tracheobronchial epithelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RHABDITIDAE; PELODERINAE; CAENORHABDITIS
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LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDATE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 332; DB 5;
Pred. No. 1.15e-12;
59; Mismatches 84
                                  Score 327; DB 6;
Pred. No. 2.69e-12;
                                                                                                                                                                                                                                                                                                                                                                                                               ; VERTEBRATA; MAMMALIA; CERCOPITHECINAE; MACACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                 10:546-551(1994)
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                                                                                                            906FCB59 CRC32;
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01-NOV-1996
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01-JUN-1998 (TREMBLREL. 06, LA
HYPOTHETICAL 170.5 KD PROTEIN.
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Q14888
Q14888;
SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).
EUKARYOTA; FUNGI; ASCOMYCOTA; ARCHIASCOMYCETES;
SCHIZOSACCHAROMYCETALES; SCHIZOSACCHAROMYCETACEAE;
                                                                                                                                                                                                                          042854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE-TRACHEOBRONCHIAL MUCOSA; GUYONNET-DUPERAT V., AUDIE J., 1 ZOUITINA-GALIEGUE S., PIGNY P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HOMO SAPIENS (HUMAN).
EUKARYOTA; METAZOA; CHORDATA;
                                                                                                                                                                                                     042854;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   430 STPSPVPPSTPLLSAHSKTGSRDCSTQTERGTESNKTAAVAPISVPAPVAAAATAAAITA 489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  113 TSAPISST-TSATTTSTTSGPGTTPSPVPTTSTTSAPTTSTT-SGPGTTPSAVPTTSITS 170
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                                                                                                                                                                                                                                                                                                                         610 TPAVAQAEVPASPATGPGPHRLSIPSLTCNPDKTDGPV
                                                                                                                                                                                                                                                                                                                                                                       230 TSTISVPTTSTTSASTTSTTSASTTSTTSGPGTTPSPV 267
                                                                                                                                                                                                                                                                                                                                                                                                                          550 VAPSAAAAAVQVAPAAPAPVPAPALVPVPAPAAAQASAPAQTQAPTSAPAVAPTPAPTP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54 TTPRPVP-TTSTTSSPTTSTTSAPTTSTTSASTTSTTSGAGTTPSPVPTTSTTSAPTTRT 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    72
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Similarity 22.08;
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279 AA;
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Pred. No. 2.27e-12;
91; Mismatches 75
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, DEGAND P., AUBERT J., P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41852EDA CRC32;
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Matches 6
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MEDLINE; 982644y.

TELFORD E.A.R., WATSON M.S., PERKY U., TELFORD E.A.R., WATSON M.S., PERKY U.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CULLINANE A.A., NEILAN J., WILSON L., DAVISON "The DNA sequence of the equine herpesvirus 4 glycoprotein gpl7/18, the homologue of herpes glycoprotein gD.";
J. GEN. VIROL. 74:0-0(0).
                                                                                                                                                                                                                                                                    NAGESHA H.S., CRABB B.S., STUDDERT M.J.; *Analysis of the nucleotide sequence of five the unique short region of the equine herpesvaRCH. VIROL. 128:143-154(1993).
                                                                                                                                                                                 STRAIN-NS80567
                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE: 93119267
                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-NS80567
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE OF 1-111 FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-NS80567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-1998 (TREMBLREL. 0:
01-JAN-1998 (TREMBLREL. 0:
01-NOV-1998 (TREMBLREL. 0:
POSITIONAL COUNTERPART OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1177
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SUBMITTED (FEB-1998) TO EN
EMBL; ALOZ1813; E1250590;
HYPOTHETICAL PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ALPHAHERPESVIRINAE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VIRUSES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EQUINE HERPESVIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MURPHY L., HARRIS D.;
SUBMITTED (FEB-1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       546
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     429
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TATAATITTTMVAAAPVAVAAAAAPAA-AAAPSPATAAATAAA-VSPAAAGQIPAAASVA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DSDNA VIRUSES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1611 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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VARICELLOVIRUS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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O EMBL/GENBANK/DDBJ
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8, LAST ANNOTATION UPDAT
HSV-1 GENE US5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CREATED)
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Pred. No. 7.45e-12;
83; Mismatches 69
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4 gene encoding
s simplex virus
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                                                                                                                  A.A.,
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                                                                                                                  A.J.;
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Best Local
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Best Local :
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BONFIELD J., BURTON J., CONNELL M., COPSEY T., COOPER J., COULSO!
CRAXTON M., DEAR S., DU Z., DURBIN R., FAVELLO A., FULTON L.,
GARDNER A., GREEN P., HAWKINS T., HILLIER L., JIER M., JOHNSTON I
JONES M., KERSHAW J., KIRSTEN J., LAISTER N., LATREILLE P.,
JONES M., KERSHAW J., KIRSTEN J., LAISTER N., LATREILLE P.,
LIGHTNING J., LLOYD C., MCMURRAY A., MORTIMORE B., O'CALLAGHAN M.
PARSONS J., PERCY C., RIFKEN L., ROOPEA, SUUNDERS D., SHOWNKEI
SMALDON N., SMITH A., SONNHAMMER E., STADEN R., SULSTON J.,
THIERRY-MIEG J., THOMAS K., VAUDIN M., VAUGHAN K., WATERSTON R.,
WANTSON A., WEINSTOCK L., WILKINSON-SPROAT J., WOHLDMAN P.;
                                                                                                                                                                                                                                                                                                                                CAENORHABDITIS ELEGANS.
FIIKARYOTA: METAZOA; NEMATODA;
                                                                                                                                                                                                                                                                                                                                                                                                                Q20908
Q20908;
                                                                                                                                                                                                                                           MEDLINE;
                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                            SUBMITTED
                                                                                                                                                                                                                                                                                                  SEQUENCE
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01-NOV-1996 (TREMBLREL.
01-JAN-1999 (TREMBLREL.
                                                                                                               WATSON A., WEINSTOCK I
                                                                                                                                                                                                                                                                                                                        RHABDITINA;
                                                                                                                                                                                                                                                                                                                                                          F56H9.1.
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 TITVPVSTTTSTSSTTTTTVPSTTTSSATTTTVPSTTTSSTTTTVPPTTTSTTTTTVP
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Similarity 27.6%;
55; Conservation
                                  Similarity 19.7%;
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                                                                                                                                                                                                                                                                                                                       RHABDITOIDEA;
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                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        624
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                                                                                                                                                                                                                                                                                                                       RHABDITIDAE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PERRY J.,
                      Score 307; DB 5;
Pred. No. 7.92e-11
95; Mismatches 6
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LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDAT
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Pred. No. 3.41e-11
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                                                                                                                                                                                                                                                                                                                                SECERNENTEA;
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                                                                 771AB928 CRC32;
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                                                                                                                                                                                                                                                                                                                      PELODERINAE;
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                                                                                                               J., WOHLDMAN from chromosc
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                                                                                                                                                                                                                                                                                                                               RHABDITIA; RHABDITIDA
                                                                                                                                                                                                                                                                                                                                                                             UPDATE)
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                        60;
                                            Length
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BANKS
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                                                                                                                                                          LAGHAN M.,
SHOWNKEEN
                                                                                                               III
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Best Local Similarity
Matches 53; Conser
  017893
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WATERSTON R.; SUBMITTED (MAR-1996) TO EMBL/GENBANK/DDBJ DATA BANKS EMBL; U51998; G1255319; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WATSON A., WEINSTOCK L., WILKINSON-SPROAT
"2.2 Mb of contiguous nucleotide sequence elegans.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WILSON R., AINSCOUGH R., ANDERSON K., BAYNES C., BERKS M.,
BORTIELD J., BURTON J., CONNELL M., COPSEY T., COOPER J., COULSON
CRAXTON M., DEAR S., DU Z., DURBIN R., FAVELLO A., FULTON L.,
GARDNER A., GREEN P., HAWKINS T., HILLIER L., JIER M., JOHNSTON L.
JONES M., KERSHAW J., KIRSTEN J., LAISTER N., LATTREILLE P.,
LIGHTWING J., LLOYD C., MCMURRAY A., MOTIMORE B., O'CALLAGHAN M.,
PARSONS J., PERCY C., RIFKEN L., ROOPRA A., SAUNDERS D., SHOWNKEED,
SMALDON N., SMITH A., SONNHAMMER E., STADEN R., SULSTON J.,
THIERRY-MIEG J., THOMAS K., VAUDIN M., VAUGHAN K., WATESTON R.,
WATESON A., WEINSTOCK L., WILKINSON-SPROAT J., WOHLDMAN P.;
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C12D12.1.
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01-NOV-1996 (TREMBLREL.
01-NOV-1996 (TREMBLREL.
01-NOV-1998 (TREMBLREL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAENORHABDITIS ELEGANS.
EUKARYOTA; METAZOA; NEMATODA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q17921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-BRISTOL N2;
NHAN M., HAWKINS J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE; 94150718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RHABDITINA;
                                                                                                                                                                                                                      542
                                                                                                                                                                                                                                                                                                                                                                     614 TIPTSQPPVTTTSLLTTLTIPTVPVTTTVVPSSATVPTTPPTTVTVAATTTSKAPVVTTS 673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             554 TTASOPTITKPVVTTNSVTPSTGTTTVPVPTTTGSPTTQTTAPVTKPTVPSSTTQTAPPV 613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBMITTED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                             601
                                                                                                                                                          734 VPPVTTTTPTTTTSTTTTTTTTT
                                                                                                                                                                                                                                                                                                                              488
                                                                                                                                                                                                                                                                                                                                                                                                                                      430
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                155
                                                                                                                                                                                                             ASVASAAAVAPSAAAAAVQVAPAAPAPVPAPALVPV-PAPAAAQASAPAQTQAPTSAPA
                                                                                                                                                                                                                                                             PTLAPTSPTKLPTSPPSTVGTSPTAPANLTTPTTAPVNPTSSTTAPTAPVNPTSPTTAPT 733
                                                                                                                                                                                                                                                                                                                   TA-TA-ATITTT-MVAA--AP-VAVAAAAAPAAAAAPSPATAAATAAAVSPAAAGQIPAA
                                                                                                                                                                                                                                                                                                                                                                                                                           STPSPVPPSTPLLSAHSKTGSRDCST-QTERGTESNKTAAVAPISVPA-PVAAAATAAAI 487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSTTTTTTTT-LPTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA-VAPSAAAAAVQVAPAAPAPVPAPALVPVPAPAAAQASAPAQTQAPTSAPAVAPTPA 606
                                                                                                        VAPTPAPTPTPAVAQAEVPASPAT 624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PTPTPAVAQAEVPASPAT 624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITVPPITTSTITTTVPATTTSTATTTTVPPTTSTITTTVPPTTTSTTTTTTVPPTTT 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (APR-1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       825 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RHABDITOIDEA;
PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRELIMINARY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.9%;
26.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86617 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TO EMBL/GENBANK/DDBJ DATA BANKS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DDA; SECERNENTEA; RHABDITIA; RHABDITIDA; RHABDITIDA; RHABDITIDAE; PELODERINAE; CAENORHABDITIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
81; M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 306; DB 5;
Pred. No. 9.37e-11
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LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDATE)
  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90FA8FCE CRC32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches 61;
  851 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 825 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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J., COULSON A.,
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Search completed: Tue Jan 18 09:51:31 2000 Job time: 47 secs.

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Best Local
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elegans ";
NATURE 368:32-38(1994).
EMBL; 283318; E1347093;
CHOTTENCE 851 AA; 9212
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BONFIELD J., BURTON J., CONNELL M., COPSEY T., COOPER J., COULSON A.
CRAXTON M., DEAR S., DU Z., DURBIN R., FAVELLO A., FULTON L.,
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JONES M., KERSHAW J., KIRSTEN J., LAISTER N., LATREILLE P.,
JONES M., KERSHAW J., KIRSTEN J., LAISTER N., LATREILLE P.,
LIGHTNING J., LLOYD C., MCMURRAY A., MORTINGE B., O'CALLAGHAN M.,
PARSONS J., PERCY C., RIFKEN L., ROOPRA A., SAUNDERS D., SHOMNKEEN F.
SMALDON N., SMITH A., SONNHAMMER E., STADEN R., SULSTON J.,
THIERRY-MIEG J., THOMAS K., VAUDIN M., VAUGHAN K., WATERSTON R.,
WATSON A., WEINSTOCK L., WILKINSON-SPROAT J., WOHLDMAN P.;
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SUBMITTED
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EUKARYOTA; METAZOA; NEMATODA; SECERNENTEA; RHABDITIA; RHABDITIDA;
RHABDITINA; RHABDIÇOIDEA; RHABDITIDAE; PELODERINAE; CAENORHABDITI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                017893;
01-JAN-1998 (TREMBLREL.
01-JAN-1998 (TREMBLREL.
01-JAN-1999 (TREMBLREL.
                                                                                                                                                                                                                                                                                                                                                                                                       WAISON A., WEINSTOCK I., WILKINSUN-SEKUAI J., WULLDELLE ... I of C. "2.2 Mb of contiguous nucleotide sequence from chromosome III of C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   F55B11.3
                                      585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE;
   602
                                                                        545
                                                                                                                                                                             465
                                                                                                                                                                                                         428 VPSTPSPVPPSTPLLSAHSKTGSRDCST-QTERGTESNKTAAVAPISVPAPVAAAATAAA 486
                                                                                                                                       487 ITATAATITTTMVAAAPVAVAAAAAAPAAAAA-PSPATAAATAAAVSPAAAGQIPAAAS-V
                                                                                                                                                                                                                                           405 VPTTTTTVPTTTTVPTTTTSVPTTTTTVPTTTTTVPTTTTTVSTTTTTVPTTTTTVPTTTTTVSTT 464
                                                                                                                                                                                                                                                                                                 Local
                                TTVPTTTTTVSTTTTTVPTTTTTVPTTTT-TVPTTTTVPTTTTPVL 630
APTPAPTPTPAVAQAEVPASPATGPGPHRLSIPSLTCNPDKTDGPVF 648
                                                                                                     PTTTTTVPTTTTTVSTTTTTVPTTTTTVPTTTTTVSTTTTTVPTTTTAPTTTTVPTTT
                                                                                                                                                                        TITVPTTTTTVPTTTTTVPTTTTTVPTTTTSVPTTTTVPTTTTVPTTTTTVSTTTTTV 524
                                                                   ASAAAVAPSAAAA-AAVQVA-PAAPAPVPAPAL-VPVPAPAAAQASAPAQTQAPTSAPAV
                                                                                                                                                                                                                                                                               h 6.9%;
Similarity 18.9%;
43; Conservative
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                                                                                                                                                                                                                                                                                                                                                      92127 MW;
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                                                                                                                                                                                                                                                                             Score 304; DB 5;
Pred. No. 1.31e-10;
94; Mismatches 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CREATED)
LAST SEQUENCE UPDATE)
LAST ANNOTATION UPDATE)
                                                                                                                                                                                                                                                                                                                                                  87CF937F CRC32;
                                                                                                                                                                                                                                                                                                                Length 851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BANKS
                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COULSON A.,
                                                                                                                                                                                                                                                                             Gaps
                                                                                                     584
                                                                                                                                       544
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US-08-403-379A-1
                          Sequence 1, Application US/08403379A
Patent No. 5756662
                           GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DETECTION
TITLE OF INVENTION: OF T. CRUZI INFECTION
NUMBER OF SEQUENCES: 9
CCRRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
(ITY Seattle
STATE: Washington
CCUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/403,379A
FILING DATE: 14-MAR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.406
TELECOMMUNICATION INFORMATION:
TELEPONE: (206) 682-6031
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 262 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-403-379A-1

6.4%; Score 213; DB 1; Len
     Othery Match
Best Local Similarity 48.4%; Score 213; DB 1; Length 262;
Matches 74; Conservative 9; Mismatches 54; Indels
               594 APTSAPAVAPTPAPTPTPAVAQAEVPASPATGP 626
             223 AAPAKAATAPAKAAT - APAKA - AAAPAKAATAP 253
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RESULT

Qy Db

Qy

Qy

Db

RESULT

32-063